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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
10/786,425	02/25/2004	Joseph A. Sorge	225436/1163	2805	
27495 PALMER & D	7590 04/11/2007 ODGE, LLP	EXAMINER			
KATHLEEN N	A. WILLIAMS / STR		KAM, CHIH MIN		
BOSTON, MA	GTON AVENUE . 02199		ART UNIT	PAPER NUMBER	
,			1656		
					
SHORTENED STATUTOR	RY PERIOD OF RESPONSE	MAIL DATE	. DELIVERY MODE		
3 MO	NTHS	04/11/2007 PAPER			

Please find below and/or attached an Office communication concerning this application or proceeding.

If NO period for reply is specified above, the maximum statutory period will apply and will expire 6 MONTHS from the mailing date of this communication.

		Applic	ation No.	Applicant(s)	
		10/78	6,425	SORGE ET AL.	
Office Action Summary			ner	Art Unit	
			lin Kam	1656	
Period fo	The MAILING DATE of this communicator Reply	ion appears on	the cover sheet	with the correspondence a	ddress
WHIC - Exte after - If NC - Failu Any	ORTENED STATUTORY PERIOD FOR CHEVER IS LONGER, FROM THE MAIL nsions of time may be available under the provisions of 31 SIX (6) MONTHS from the mailing date of this communic or period for reply is specified above, the maximum statutor re to reply within the set or extended period for reply will, reply received by the Office later than three months after the patent term adjustment. See 37 CFR 1.704(b).	ING DATE OF CFR 1.136(a). In nation. The period will apply as the statute, cause the	THIS COMMUN o event, however, may nd will expire SIX (6) Mo application to become	NICATION. a reply be timely filed ONTHS from the mailing date of this ABANDONED (35 U.S.C. § 133).	
Status			-	•	
1) 又	Responsive to communication(s) filed o	n <i>22 January 2</i>	2007.		
-	_	This action			
3)□	Since this application is in condition for	allowance exc	ept for formal ma	atters, prosecution as to th	ne merits is
	closed in accordance with the practice u	ınder <i>Ex par</i> te	Quayle, 1935 C	.D. 11, 453 O.G. 213.	
Disposit	on of Claims				
4)⊠	Claim(s) 1-10 is/are pending in the appl	ication.			
	4a) Of the above claim(s) is/are v		consideration.		•
5)⊠	Claim(s) 1 is/are allowed.			•	
6)⊠	Claim(s) 2-10 is/are rejected.		٠		
7)	Claim(s) is/are objected to.				
8)[Claim(s) are subject to restriction	and/or election	n requirement.		
Applicat	on Papers				
9)[The specification is objected to by the Ex	kaminer.	•		
10)⊠	The drawing(s) filed on <u>25 February 200</u>	<u>4</u> is/are: a)⊠	accepted or b)	objected to by the Exam	niner.
	Applicant may not request that any objection	to the drawing	s) be held in abey	ance. See 37 CFR 1.85(a).	
	Replacement drawing sheet(s) including the	correction is red	quired if the drawir	ng(s) is objected to. See 37 (CFR 1.121(d).
11)	The oath or declaration is objected to by	the Examiner.	Note the attach	ed Office Action or form P	PTO-152.
Priority ι	ınder 35 U.S.C. § 119				
	Acknowledgment is made of a claim for t ☐ All b) ☐ Some * c) ☐ None of:	foreign priority	under 35 U.S.C.	§ 119(a)-(d) or (f).	
	1. Certified copies of the priority documents have been received.				
	2. Certified copies of the priority doc				
	3. Copies of the certified copies of the			en received in this Nationa	l Stage
* ~	application from the International	•	` ''		
* 8	see the attached detailed Office action fo	r a list of the c	ertified copies no	ot received.	
				•	
Attachmen	t(s)				
	e of References Cited (PTO-892)	240		/ Summary (PTO-413)	
	e of Draftsperson's Patent Drawing Review (PTO-s nation Disclosure Statement(s) (PTO/SB/08)	/4 8)		o(s)/Mail Date f Informal Patent Application	
Paper No(s)/Mail Date 6) Other: <u>sequence aligment</u> .					

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DETAILED ACTION

Status of the Claims

1. Claims 1-10 are pending.

Applicants' amendment filed January 22, 2007 is acknowledged. Applicant's response has been fully considered. Claims 1-10 have been amended. Therefore, claims 1-10 are examined.

Withdrawn Informalities

2. The previous objection to the specification, regarding certain sequences not having sequence identifier, is withdrawn in view of applicants' amendment to the specification in the amendment filed January 22, 2007.

Withdrawn Claim Objection

3. The previous objection to claims 1 and 5, is withdrawn in view of applicants' amendment to the claim, and applicants' response at page 5 in the amendment filed January 22, 2007.

Withdrawn Claim Rejections - 35 USC § 112

4. The previous rejection of claims 2-4 and 6-10 under 35 U. S. C. 112, second paragraph, is withdrawn in view of applicants' amendment to the claims, and applicants' response at page 5 in the amendment filed January 22, 2007.

Withdrawn Claim Rejections - 35 USC § 103

5. The previous rejection of claims 1 and 3-5 under 35 U. S. C. 103(a) as being unpatentable over Zolotukhin *et al.* (U.S. Patent 5,874,304) in view of Bryan *et al.* (U. S. Patent 6,232,107), and applicants' response at pages 5-8 in the amendment filed January 22, 2007.

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New Claim Rejections - 35 USC § 112

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

6. Claims 2-4 and 6 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

Claims 2-4 and 6 are directed to a humanized polynucleotide, the polynucleotide encoding R. reniformis GFP having the sequence of SEQ ID NO:2, wherein the polynucleotide comprises SEQ ID NO:3; a recombinant vector comprising the humanized polynucleotide; a cell comprising the recombinant vector; and a method of producing R. reniformis GFP having the sequence of SEQ ID NO:2 using the humanized polynucleotide. While the specification indicates the present invention encompasses recombinant polynucleotides encoding the GFP from R. reniformis, where the polynucleotide sequence is a humanized sequence such as SEQ ID NO:3 (page 5, lines 8-10; page 6, line 23; Fig. 5; page 11, line 14-page 13, line 4; page 17, lines 9-10; Table 1 at page 50), the specification does not disclose a humanized polynucleotide encoding R. reniformis GFP having the sequence of SEQ ID NO:2 comprises the sequence of SEQ ID NO:3. A sequence alignment of a nucleotide sequence encoding SEQ ID NO:2 with SEQ ID NO:3 indicates SEQ ID NO:3 does not encode SEQ ID NO:2 (see attached sequence alignments; Fig 5), e.g., SEQ ID NO:2 has Val at position 56, Ser at position 131, and Asp at

position 208, while SEQ ID NO:3 has the corresponding nucleotides encoding amino acid at position 56 are GCC (which encodes Ala), nucleotides encoding amino acid at position 131 are CGC (which encodes Arg, although Fig. 5 shows Ser at this position); and nucleotides encoding amino acid at position 208 are TAC (which encodes Tyr). The lack of description on the humanized polynucleotide encoding R. reniformis GFP having the sequence of SEQ ID NO:2 comprises SEQ ID NO:3 and lack of representative species as encompassed by the claims, applicants have failed to sufficiently describe the claimed invention, in such full, clear, concise terms that a skilled artisan would not recognize applicants were in possession of the claimed invention.

New Claim Rejections - 35 USC § 112

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

7. Claims 5 and 7-10 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

Claims 5 and 7-10 are indefinite because the claim recites a humanized polynucleotide sequence encoding R. reniformis GFP in step (a) without identifying the sequence of R. reniformis GFP and providing "SEQ ID NO:", thus it is not clear what sequence the R. reniformis GFP has. Claims 7-10 are included in the rejection because they are dependent on a rejected claim and do not correct the deficiency of the claim from which they depend.

Conclusion

8. Claims 2-10 are rejected; and it appears that claim 1 is free of art.

Applicant's amendment necessitated the new ground(s) of rejection presented in this Office action. Accordingly, **THIS ACTION IS MADE FINAL**. See MPEP § 706.07(a). Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for reply to this final action is set to expire THREE MONTHS from the mailing date of this action. In the event a first reply is filed within TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the date of this final action.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Chih-Min Kam whose telephone number is (571) 272-0948. The examiner can normally be reached on 8.00-4:30, Mon-Fri.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Kathleen Bragdon can be reached at 571-272-0931. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Chih-Min Kam, Ph. D.

Primary Patent Examiner

CHIH-MIN KAM PRIMARY EXAMINER

CMK

April 9, 2007

Sequence Alignment of SERSON: 2 against nucleotide sequence

ALIGNMENTS

RESULT 1

```
US-10-786-425-1
; Sequence 1, Application US/10786425
 GENERAL INFORMATION:
  APPLICANT: Stratagene
  TITLE OF INVENTION: Renilla Reniformis Green Fluorescent Protein and Mutants
  FILE REFERENCE: 25436/1162
  CURRENT APPLICATION NUMBER: US/10/786,425
  CURRENT FILING DATE: 2004-02-25
  PRIOR APPLICATION NUMBER: US/09/795,040
  PRIOR FILING DATE: 2001-02-26
  PRIOR APPLICATION NUMBER: US 60/185,589
  PRIOR FILING DATE: 2000-02-28
  PRIOR APPLICATION NUMBER: 60/210,561
  PRIOR FILING DATE: 2000-06-09
  NUMBER OF SEQ ID NOS: 8
  SOFTWARE: PatentIn version 3.0
 SEO ID NO 1
   LENGTH: 720
   TYPE: DNA
   ORGANISM: Renilla reniformis
   FEATURE:
   NAME/KEY: exon
   LOCATION: (1)..(720)
   OTHER INFORMATION: open reading frame
US-10-786-425-1
Alignment Scores:
Pred. No.:
                    1.52e-131
                                 Length:
                                              720
Score:
                    1253.00
                                 Matches:
                                              239
Percent Similarity:
                    100.0%
                                 Conservative:
                                              0
Best Local Similarity:
                    100.0%
                                 Mismatches:
                                              0
Query Match:
                    100.0%
                                 Indels:
                                              0
DB:
                                 Gaps:
                                              0
US-10-786-425-2 (1-239) x US-10-786-425-1 (1-720)
Qу
          1 MetValSerLysGlnIleLeuLysAsnThrGlyLeuGlnGluIleMetSerPheLysVal 20
            Db
          1 ATGGTGAGTAAACAAATATTGAAGAACACTGGATTGCAGGAGATCATGTCGTTTAAAGTG 60
         21 AsnLeuGluGlyValValAsnAsnHisValPheThrMetGluGlyCysGlyLysGlyAsn 40
Qу
            Db
         61 AATCTGGAAGGTGTAGTAAACAATCATGTGTTCACAATGGAAGGTTGTGGAAAAGGAAAT 120
Qу
         41 IleLeuPheGlyAsnGlnLeuValGlnIleArgValThrLysGlyValProLeuProPhe 60
            Db
        121 ATTTTATTCGGAAACCAACTGGTTCAGATTCGTGTCACAAAAGGGGTCCCGCTTCCATTT 180
         61 AlaPheAspIleLeuSerProAlaPheGlnTyrGlyAsnArgThrPheThrLysTyrPro 80
Qy
            Db
        181 GCATTTGATATTCTCTCACCAGCTTTCCAATACGGCAACCGTACATTCACGAAATACCCG 240
         81 GluAspIleSerAspPhePheIleGlnSerPheProAlaGlyPheValTyrGluArqThr 100
Qy
            Db
        241 GAGGATATATCAGACTTTTTTATACAATCATTTCCAGCGGGATTTGTATACGAAAGAACG 300
        101 LeuArgTyrGluAspGlyGlyLeuValGluIleArgSerAspIleAsnLeuIleGluGlu 120
Qy
            Db
        301 TTGCGTTACGAAGATGGTGGACTGGTTGAAATCCGTTCAGATATAAATTTAATCGAGGAG 360
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121 MetPheValTyrArqValGluTyrLysGlySerAsnPheProAsnAspGlyProValMet 140
Qy
            361 ATGTTTGTCTACAGAGTGGAATATAAAGGTAGTAACTTCCCGAATGATGGTCCAGTGATG 420
Db
        141 LysLysThrIleThrGlyLeuGlnProSerPheGluValValTyrMetAsnAspGlyVal 160
Qy
            421 AAGAAGACAATCACAGGATTACAACCTTCGTTCGAAGTTGTGTATATGAACGATGGCGTC 480
Db
        161 LeuValGlyGlnValIleLeuValTyrArgLeuAsnSerGlyLysPheTyrSerCysHis 180
Qy
            481 TTGGTTGGCCAAGTCATTCTTGTTTATAGATTAAACTCTGGCAAATTTTATTCGTGTCAC 540
Db
        181 MetArgThrLeuMetLysSerLysGlyValValLysAspPheProGluTyrHisPheIle 200
Qy
            541 ATGAGAACACTGATGAAATCAAAGGGTGTAGTGAAGGATTTTCCCGAATACCATTTCATT 600
Db
        201 GlnHisArgLeuGluLysThrAspValGluAspGlyGlyPheValGluGlnHisGluThr 220
Qу
            601 CAACATCGTTTAGAGAAGACTGATGTGGAAGACGGAGGTTTTGTTGAGCAACACGAGACG 660
Db
Qy
        221 AlaIleAlaGlnLeuThrSerLeuGlyLysProLeuGlySerLeuHisGluTrpVal 239
            661 GCCATTGCTCAACTGACATCGCTGGGGAAACCACTTGGATCCTTACACGAATGGGTT 717
Db
RESULT 2
US-10-786-425-3
; Sequence 3, Application US/10786425
; GENERAL INFORMATION:
  APPLICANT: Stratagene
  TITLE OF INVENTION: Renilla Reniformis Green Fluorescent Protein and Mutants
  FILE REFERENCE: 25436/1162
  CURRENT APPLICATION NUMBER: US/10/786,425
  CURRENT FILING DATE: 2004-02-25
  PRIOR APPLICATION NUMBER: US/09/795,040
  PRIOR FILING DATE: 2001-02-26
  PRIOR APPLICATION NUMBER: US 60/185,589
  PRIOR FILING DATE: 2000-02-28
  PRIOR APPLICATION NUMBER: 60/210,561
  PRIOR FILING DATE: 2000-06-09
  NUMBER OF SEQ ID NOS: 8
  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 3
   LENGTH: 720
   TYPE: DNA
   ORGANISM: Artificial Sequence
   FEATURE:
   OTHER INFORMATION: Humanized DNA
   FEATURE:
   NAME/KEY: exon
   LOCATION: (1)..(720)
   OTHER INFORMATION: open reading frame
US-10-786-425-3
Alignment Scores:
Pred. No.:
                    1.64e-129
                                 Length:
                                              720
                   1235.00
                                 Matches:
Score:
                                             236
Percent Similarity:
                    98.7%
                                 Conservative:
                                             0
Best Local Similarity:
                    98.7%
                                 Mismatches:
                                             3
```

Indels:

٥

Query Match:

98.6%

DB: 7 Gaps: 0

US-10-786-425-2 (1-239) x US-10-786-425-3 (1-720)

Qy	1	MetValSerLysGlnIleLeuLysAsnThrGlyLeuGlnGluIleMetSerPheLysVal 20
Db	1	ATGGTGAGCAAGCAGATCCTGAAGAACACCGGCCTGCAGGAGATCATGAGCTTCAAGGTG 60
Qy	21	AsnLeuGluGlyValValAsnAsnHisValPheThrMetGluGlyCysGlyLysGlyAsn 40
Db	61	AACCTGGAGGGCGTGGTGAACAACCACGTGTTCACCATGGAGGGCTGCGGCAAGGGCAAC 120
Qy	41	<pre>IleLeuPheGlyAsnGlnLeuValGlnIleArgValThrLysGlyValProLeuProPhe 60 </pre>
Db	121	ATCCTGTTCGGCAACCAGCTGGTGCAGATCCGCGTGACCAAGGGCGCCCCCCTGCCCTTC 180
Qy	61	AlaPheAspIleLeuSerProAlaPheGlnTyrGlyAsnArgThrPheThrLysTyrPro .80
Db	181	GCCTTCGACATCCTGAGCCCCGCCTTCCAGTACGGCAACCGCACCTTCACCAAGTACCCC 240
Qy	81	GluAspIleSerAspPhePheIleGlnSerPheProAlaGlyPheValTyrGluArgThr 100
Db	241	GAGGACATCAGCGACTTCTTCATCCAGAGCTTCCCCGCCGGCTTCGTGTACGAGCGCACC 300
Qy	101	LeuArgTyrGluAspGlyGlyLeuValGluIleArgSerAspIleAsnLeuIleGluGlu 120
Db	301	CTGCGCTACGAGGACGCCGGCCTGGTGGAGATCCGCAGCGACATCAACCTGATCGAGGAG 360
Qy	121	MetPheValTyrArgValGluTyrLysGlySerAsnPheProAsnAspGlyProValMet 140
Db		ATGTTCGTGTACCGCGTGGAGTACAAGGGCCGCAACTTCCCCAACGACGGCCCCGTGATG 420
Qy	141	LysLysThrIleThrGlyLeuGlnProSerPheGluValValTyrMetAsnAspGlyVal 160
Db	421	AAGAAGACCATCACCGGCCTGCAGCCCAGCTTCGAGGTGGTGTACATGAACGACGGCGTG 480
Qy	161	LeuValGlyGlnValIleLeuValTyrArgLeuAsnSerGlyLysPheTyrSerCysHis 180
Db	481	CTGGTGGCCAGGTGATCCTGGTGTACCGCCTGAACAGCGGCAAGTTCTACAGCTGCCAC 540
Qy	181	MetArgThrLeuMetLysSerLysGlyValValLysAspPheProGluTyrHisPheIle 200
Db	541	ATGCGCACCTGATGAAGAGCAAGGGCGTGGTGAAGGACTTCCCCGAGTACCACTTCATC 600
Qy	201	GlnHisArgLeuGluLysThrAspValGluAspGlyGlyPheValGluGlnHisGluThr 220
Db	601	CAGCACCGCCTGGAGAAGACCTACGTGGAGGACGGCGGCTTCGTGGAGCACGAGACC 660
Qy	221	AlaIleAlaGlnLeuThrSerLeuGlyLysProLeuGlySerLeuHisGluTrpVal 239
Db ·	661	GCCATCGCCCAGCTGACCAGCCTGGGCAAGCCCCTGGGCAGCCTGCACGAGTGGGTG 717